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Minimum
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Perfect score:
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Copyright (c) 1993 - 2000 Comp
                    sp_plant:*
sp_rodent:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed. is derived by analysis of the total score distribution.

SUMMARIES

Result

		de C				
Result No.	Score	Query Match	Length	DB	ID	Description
1	384	100.0	966	10	Q9SPL5	Q9spl5 macadamia i
ν	367	95.6	666	10	Q9SPL4	
ω	363	94.5	625	10	Q9SPL3	
4	172	44.8	525	10	Q43358	Q43358 theobroma c
ъ	163	42.4	593	10	Q9SEW4	
6	120.5	31.4	554	ഗ	Q9VPS3	Q9vps3 drosophila
7	116.5	30.3	1038	Ŋ	060983	060983 dictyosteli
8	115	29.9	1089	12	040947	040947 kaposi's sa
9	114	29.7	810	10	Q9ZWI3	cucurbita
10	107	27.9	1162	12	Q98148	
11	105.5	27.5		12	Q9QR71	kaposi's
12	105	27.3	1737	ഗ	Q9TW28	dictyostel
13	104.5	27.2	385	ഗ	Q17909	Q17909 caenorhabdi
14	104	27.1	1390	S	077033	
15	103	26.8	838	տ	Q9Y0C9	Q9y0c9 dictyosteli
16	101	26.3	233	4	Q9UFP2	
17	101	26.3	289	4	Q9NUN6	
18	101	26.3	1027	4	Q9NST7	homo
19	101	26.3	1165	4	095819	homo

RESULT Q9SPL4

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Q9UNV6 nomo sapien Q61118 mus musculu	homo	_			Q9w3x0 drosophila	Q25777 plasmodium	Q9wu62 mus musculu	Q9txb8 dictyosteli	Q9vva9 drosophila	Q9uke5 homo sapien			homo	homo					Q9y520 homo sapien	Q9u9s7 dictyosteli	Q9sp11 glycine max	O95033 homo sapien		075172 homo sapien

ALIGNMENTS

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                                                                                                 Query Match
Best Local S
Matches 69
                                                                                                                                                               Marcus J.P., Goulter K.C., Green J.L., Manners J.M.;
Ma family of antimicrobial peptides is produced by pi
globulin protein in Macadamia integrifolia kernels.",
Plant J. 0:0-0(1999).
EMBL; AF161883; AAD54244.1; -.
HSSP; P02853; 2PHL.
INTERPRO; IPR001113; -.
PFAM; PF00546; Seedstore_7s; 1.
SEQUENCE 666 AA; 78217 MW; C752B884B2DF0224 CRC64
                                                                                                                                                                                                                                                                                                                                                                                   OSPL5 PRELIMINARY; PRT;
OSPNT5;
O1-MAY-2000 (TrEMBLrel. 13, Created)
O1-MAY-2000 (TrEMBLrel. 13, Last sequ
O1-OCT-2000 (TrEMBLrel. 15, Last anno
VICILIN PRECURSOR.
                                                                                                                                                                                                                                                                                                                    Macadamia integrifolia (Macadamia nut).
Eukaryota; Viridiplantae; Embryophyta; Tracheophyta;
Magnoliophyta; eudicotyledons; Proteaceae; Macadamia.
NCBI_TaxID=80698;
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177
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                                              NRQRDPQQQYEQCQKHCQRRETEPRHMQTCQQRCERRYEKEKRKQQKRYEEQQREDEEKY 60
EERMKEEDN 185
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Pred. No. 5.2e-27;
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                                                                     PFAM; PF00546; Seedstore_7s; 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  globulin protein in Macadamia integrifolia.";
Plant J. 0:0-0(1999).
EMBL: AF161884; AAD54245.1;
INTERPRO: TECHNOLOGY.
                                                                                                            globulin protein in Macadamia integrifolia.";
Plant J. 0:0-0(1999).
EMBL, AF161885; AAD54246.1; -.
HSSP; P02853; 2PHL.
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Q9SPL4;
01-MAY-2000
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Macadamia integrifolia (Macadamia nut).
Eukaryota; Viridiplantae; Embryophyta; Tracheophyta;
Magnoliophyta; eudicotyledons; Proteaceae; Macadamia
                                                                                                                                              TISSUE=NUT KERNEL;
Marcus J.P., Goulter K.C.,
"A family of antimicrobial
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PFAM; PF00546; Seedston
SEQUENCE 666 AA; 782
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Q9SEW4;
01-MAY-2000
01-MAY-2000
01-OCT-2000
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-CT-2000 (TrEMBLrel. 15, Last annotation update)
VICILIN-LIKE PROTEIN PRECURSOR (FRAGMENT).
Juglans regia (English walnut).
Eukaryota; Viridiplantae; Embryophyta; Tracheophyt
Magnoliophyta; eudicotyledons; core eudicots; Rosi
Fagales; Juglandaceae; Juglans.
NCBL_TaxID-51240;
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Q1-NOV-1996
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MEDLIND=92288309; PubMed=1600151;

MCHenry L., Fritz P.J.;

Comparison of the structure and nucleotide sequences of vicilin of cocoa and cotton raise questions about vicilin evolution.";

Plant Mol. Biol. 18:1173-1176(1992).

EMBL; X62625; CAA44493.1; -.

EMBL; X62625; CAA44494.1; -.

MENDEL; X62625; CAA44494.1; -.

HSSP; P02853; 2PHL.

MENDEL; 30919; Thecc;1188;30919.
STRAIN-CV. SUNLAND; TISSUE-SOMATIC EMBRYO LINE; Teuber S.S., Jarvis K.C., Peterson W.R., Dandekar A.M., Ansari A.A.; Teuber S.S., Jarvis K.C., Peterson W.R., Dandekar A.M., Ansari A.A.; "Identification and cloning of a cDNA encoding a vicilin-like protei Jug r 2, from English walnut kernel (Juglans regia): a major food
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Eukaryota; Viridiplantae; Embr.
Magnoliophyta; eudicotyledons;
Malvales; Malvaceae; Theobroma
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Pred. No. 3.1e
20; Mismatches
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RESULT OF THE PROPERTY OF THE 
RA Adams M.D. (Celniker S.E., Holt R.A., Evans C.A., Gocayne J.D., RA Adams M.D., Celniker S.E., Holt R.A., Hoskins R.A., Galle R.F., RA Adams M.D., Celniker S.E., Li P.W., Hoskins R.A., Galle R.F., RA Adams M.D., Celniker S.E., Richards S., Ashburner M., Henderson S.N., RA George R.A., Lewis S.E., Richards S., Ashburner M., Henderson S.N., RA George R.A., Lewis S.E., Richards S., Ashburner M., Henderson S.N., RA George R.A., Champe M., Pfeiffer B.D., RA Barandon R.C., Rogers Y.H.C., Blazej R.G., Champe M., Pfeiffer B.D., RA Barandon R.C., Baxter E.G., Helt G., Nelson C.R., Miklos G.L.G., RA Baril J.F., Agbayani A., An H.-J., Andrews Pfannkoch C., Baldwin D., RA Ballew R.M., Basu A., Baxendale J., Bayraktaroglu L., Beasley E.M., RA Ballew R.M., Basu A., Baxendale J., Bayraktaroglu L., Beasley E.M., RA Beeson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S., Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davles P., RA Burtis K.C., Busam D.A., Butler H., Cadicu E., Center A., Chandra I., RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davles P., RA de Pablos B., Delcher A., Deng Z., Mays A.D., Dew II, Dietz S.M., P. RA Durbin K.J., Evangelista C.C., Ferraz C., Ferriera S., Fleischmann W., RA Glodek A., Gong F., Gorrell J.H., Gu Z., Glabart W.M., Glasser K., RA Hortin N.J., Harvey D., Helman T.J., Hernandez J.R., Houck J., Harris M., Harris N.L., Harvey D., Helman T.J., Wei M.-H., Ibegwam C., RA Harris N.L., Harvey D., Helman T.J., Wei M.-H., Ibegwam C., RA Harris N.L., Kalush F., Karpen G.H., Ke Z., Kennison J.A., Ketchum K.A., RA Harris N., Milshina N.V., Mobarry C., Morris J., Moshrefi A., RA Mount S.M., Milshina N.V., Mobarry C., Morris J., Moshrefi A., RA Mount S.M., Milshina N.V., Mobarry C., Morris J., Moshrefi A., RA Reinert K., Remington K.S., Saunders R.D.C., Scheeler F., Shen H., RA Shue B.C., Siden K.A., Nixon K., Nusskern D.R., Pacleb J.M., Resembed M.G., Shen H., Wang A.C., Stapleton M., Skupski M.P., Smith T., Shen H., Mang Sarman D.A., Weinstock G.M., Weissenbach J.,
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Best Local :
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Submitted (MAY-1998)
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Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
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(TrEMBLrel. 13, Last sequence update)
(TremBLrel. 14, Last annotation update)
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Pred. No. 2.1e-07;
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SEQUENCE
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EMBL; AE003588: AAF51469.1; -.
FLYBASE; FB9G0031273; CG2839.
INTERPRO; IPR001304; -.
PPAM: PF00059; lectim_C; 1
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Hughes J.E., Welker D.L.;
Submitted (APR-1997) to the
EMBL; AF000580; AAC14390.1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE-9819836; PubMed-9539429; Rieben W.K. Jr., Gonzales C.M., Gonza Kiyosawa H., Hughes J.E., Welker D.L.
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01-JAN-1998
01-JAN-1998
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"Multiple functional proteins are produced by cleaving Asn-Gln
of a single precursor by vacuolar processing enzyme.";
J. Biol. Chem. 274:2563-2570(1999).
                                                                                                                                                                                                                                                       Cucurbita maxima (Pumpkin) (Winter squash).

Eukaryota; Viridiplantae; Embryophyta; Tracheophyta; Spermatophyta;

Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids I;

Cucurbitales; Cucurbitaceae; Cucurbita.
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EMBL; U93872; AABB2657.1; -.
INTERPRO; IPRO02017. -
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HSSP; P02853; 2PHL.
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MEDLINE=97296220; PubMed=9151804;
Neipel F., Albrecht J.C., Flecken
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Viruses; dsDNA viruses, no RNA stage; Herpesviridae;
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MEDLINE=99107919; PubMed=9891029;
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01-JUN-2000
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MEDLINE=97048116; PubMed=8892957;
Cesarman E., Nador R.G., Bai F., I
Moore P.S., Chang Y., Knowles D.M
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Viruses; dsDNA viruses, no RNA stage; He
Gammaherpesvirinae; Rhadinovirus.
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Parry J.P.,
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Science 274:173
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"Molecular mimicry of human
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Moore P.S., Bashoff C., Weiss R.A.,
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                                                                                                  89
                                                                                                                                                                                                                                                                                                                                AA;
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و :
                                                                                                                                                                                                                                             27.9%;
                                                                                                                                                                                                                                                                                                                                135213 MW;
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02,
14,
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                                                                                                                                                                                                                         25;
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Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Kaposi sarcoma-associated
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Moore P.
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Moore P.
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Moore P.S.;
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.S.;
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Best Local S
Matches 22
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01-MAY-2000
01-MAY-2000
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01-MAY-2000
01-JUN-2000
                     EMBL; AF09
HSSP; P087
INTERPRO;
INTERPRO;
                                                                 Dictyostelium discoideum.";
Submitted (SEP-1998) to the EMBL/GenBank/DDBJ databases
EMBL; AB017910; BAAB4604.1; -
EMBL; AF090533; AAD47903.1; -
HSSP; P08799; IMND.
                                                                                                                                                                                                                                                                                                     Oishi N., Adachi H., Sutoh K.; "Dictyostelium discoideum gene for MyoM."; Submitted (SEP-1998) to the EMBL/GenBank/E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Dictyostelium discoideum (Slime mold) Eukaryota; Dictyosteliida; Dictyostel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Q9TW28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Herpesvirus Encoding a Protein Proteins 1 and 2A of Epstein Proteins 1 and 2A of Epstein J. Virol. 73:6953-6963(1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Kaposi's sarcoma-associated herpesvirus (KSHV) (Human herpesvirus Viruses; dsDNA viruses, no RNA stage; Herpesviridae; Gammaherpesvirinae; Rhadinovirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORF 73.
                                                                                                                                                                                   Geissler H., Schwarz E.C., Soldati T.;
"Identification of two novel and highly divergent myosins
                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                           STRAIN-AX2;
                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID=44689;
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PRINTS; PR01217; PRICHEXTENSN.
SEQUENCE 1129 AA; 131346 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL; AF148805; AAD46501.1; INTERPRO; IPR000533; -. INTERPRO; IPR002017; -. INTERPRO; IPR002965; -.
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                     IPR000219;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   , Aurade F., Davison A., Schulz T.F.;
Spliced Gene from Kaposi's Sarcoma-Associated
a Protein with Similarities to Latent Membrane
Epstein-Barr Virus.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 27.5%;
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Last annotation updat
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Last sequence up
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Q17909;
Q1-NOV-1996
Q1-NOV-1996
Q1-JUN-2000
                                                                                                                                                                                                                                   Wilson R., Ainscough R., Anderson K., Baynes C., Berks M.,
Bonfield J., Burton J., Connell M., Copsey T., Cooper J., Couls
Craxton M., Dear S., Du 2., Durbin R., Favello A., Fulton L.,
Gardner A., Green P., Hawkins T., Hillier L., Jier M., Johnston
Jones M., Kershaw J., Kirsten J., Laister N., Latreille P.,
Lightning J., Lloyd C., Mcmurray A., Mortimore B., O'Callaghan
Parsons J., Percy C., Rifken L., Ropra A., Saunders D., Shownk
Smaldon N., Smith A., Sonnhammer E., Staden R., Sulston J.,
Thierry-Mieg J., Thomas K., Vaudin M., Vaughan K., Waterston R.
Watson A., Weinstock L., Wilkinson-Sproat J., Wohldman P.;
"2.2 Mb of contiguous nucleotide sequence from Chromosome III o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C11G6.3 PROTEIN.
C11G6.3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Caenorhabditis elegans
                                                                                                                                                                   SEQUENCE
                                                                                                                                                                                                     Nature 368:32-38(1994).
EMBL; 270204; CAA94113.1;
                                                                                                                                                                                                                                 elegans.
                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A. MEDLINE=94150718; PubMed=7906398;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa;
Rhabditidae; Peloder
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRINTS; PR00193; MYOSINHEAVY.
PROSITE; PS00152; ATPASE_ALPHA_BETA; UNKNOWN_1
PROSITE; PS50003; PH_DOMAIN; 1.
SEQUENCE 1737 AA; 195860 MW; 39CBF9C08EFEF
                                                                                                                                                                              PFAM; PF00628;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.
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25; Conservative
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6 (TrEMBLrel. 01,
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                                                                                                   Conservative
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                                                                                                                                                                 46052 MW;
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                                                                                                Score 104.5; DB Pred. No. 0.024; 4; Mismatches
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INTERPRO; IPR001440; -.

PFAM; PF00515; TPR; 9.

SEQUENCE 1390 AA; 160421 MW; BAAB99286
                                                                                                                                               SEQUENCE FROM N.A.

SEQUENCE FROM N.A.

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CEPOIFENCE 838 AA; 94059 MW; EAAD911294226808 CRC64;
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O77033;
O1-NOV-1998 (TrEMBLrel. 08, Created)
O1-NOV-1998 (TrEMBLrel. 08, Last sequence update)
O1-JUN-2000 (TrEMBLrel. 14, Last annotation update)
                                                                                                                                                                                                                                                                                                                         O9YOC9 PRELIMINARY; PRT; 838 AA.
(9YOC9;
(01-NOV-1999 (TrEMBLrel. 12, Created)
(01-NOV-1999 (TrEMBLrel. 12, Last sequence update)
(01-NOV-1999 (TrEMBLrel. 12, Last annotation update)
RAS INTERACTING PROTEIN RIPA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Dictyostelium discoideum (Slime mold). Eukaryota; Dictyosteliida; Dictyostelium. NCBI_TaxID=44689;
                                                                                                                                                                                                                                                                                       Dictyostelium discoideum (Slime mold).
Eukaryota; Dictyosteliida; Dictyostelium
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148
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160421 MW;
                                                                                                   26.8%; Score 103; DB 5; Length 838; 23.9%; Pred. NO. 0.061; Live 33; Mismatches 18; Indels
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23.9%; Pred. No. 0.
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